



INVITED SPEAKER PRESENTATION

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Epigenetics in rheumatic diseases

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From 21st European Pediatric Rheumatology (PReS) Congress
Belgrade, Serbia. 17-21 September 2014

With enormous speed novel data are emerging about regulating the expression of the genetically encoded information [1]. This highly complex regulatory network called *epigenetics* includes acetylation, methylation, phosphorylation, sumoylation and non-coding RNAs (ncRNA), such as miRNA and lncRNAs. Our laboratory is addressing over the past decade inflammatory rheumatic diseases [2], like rheumatoid arthritis (RA), AS, SSs and pulmonary hypertension and thereby searching for the regulation of pro-inflammatory cytokines [3,4], novel diagnostic signatures and new therapeutic targets. In this regard, DNA demethylation of RA synovial cells can be modulated by targeting specific enzymes [5]. Also, miRNA signatures for new response markers are in development [6].

doi:10.1186/1546-0096-12-S1-I12

Cite this article as: Gay: Epigenetics in rheumatic diseases. *Pediatric Rheumatology* 2014 **12**(Suppl 1):112.

Disclosure of interest

None declared.

Published: 17 September 2014

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